



# Sequence Listing

<110> CHUNTHARAPAI, ANAN  
GREWAL, IQBAL  
KIM, KYUNG JIN  
YAN, MINHONG

<120> TACI Antibodies and Uses Thereof

<130> P1942R1

<140> US 10/626,914

<141> 2003-07-25

<150> US 60/398,530

<151> 2002-07-25

<160> 17

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<212> DNA

<213> Homo sapien

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 Met Arg Ser Cys Pro Glu Glu Gln Tyr Trp Asp Pro Leu Leu Gly  
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 Tyr Gln Gly Leu Glu His Arg Gly Ser Glu Ala Ser Pro Ala Leu  
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 Pro Gly Leu Lys Leu Ser Ala Asp Gln Val Ala Leu Val Tyr Ser  
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 Thr Leu Gly Leu Cys Leu Cys Ala Val Leu Cys Cys Phe Leu Val  
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Cys	Pro	His	Ile	Pro	Asp	Ser	Gly	Leu	Gly	Ile	Val	Cys	Val	Pro	
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<213> Homo sapien

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 tcagcctcag gagaattagg ttccagtccc tctaggaagc ctctgacc 1348

<210> 12  
 <211> 250  
 <212> PRT  
 <213> Homo sapien

<400> 12  
 Met Pro Ala Ser Ser Pro Phe Leu Leu Ala Pro Lys Gly Pro Pro  
 1 5 10 15  
 Gly Asn Met Gly Gly Pro Val Arg Glu Pro Ala Leu Ser Val Ala  
 20 25 30  
 Leu Trp Leu Ser Trp Gly Ala Ala Leu Gly Ala Val Ala Cys Ala  
 35 40 45  
 Met Ala Leu Leu Thr Gln Gln Thr Glu Leu Gln Ser Leu Arg Arg  
 50 55 60

Glu	Val	Ser	Arg	Leu	Gln	Gly	Thr	Gly	Gly	Pro	Ser	Gln	Asn	Gly	65	70	75
Glu	Gly	Tyr	Pro	Trp	Gln	Ser	Leu	Pro	Glu	Gln	Ser	Ser	Asp	Ala	80	85	90
Leu	Glu	Ala	Trp	Glu	Asn	Gly	Glu	Arg	Ser	Arg	Lys	Arg	Arg	Ala	95	100	105
Val	Leu	Thr	Gln	Lys	Gln	Lys	Lys	Gln	His	Ser	Val	Leu	His	Leu	110	115	120
Val	Pro	Ile	Asn	Ala	Thr	Ser	Lys	Asp	Asp	Ser	Asp	Val	Thr	Glu	125	130	135
Val	Met	Trp	Gln	Pro	Ala	Leu	Arg	Arg	Gly	Arg	Gly	Leu	Gln	Ala	140	145	150
Gln	Gly	Tyr	Gly	Val	Arg	Ile	Gln	Asp	Ala	Gly	Val	Tyr	Leu	Leu	155	160	165
Tyr	Ser	Gln	Val	Leu	Phe	Gln	Asp	Val	Thr	Phe	Thr	Met	Gly	Gln	170	175	180
Val	Val	Ser	Arg	Glu	Gly	Gln	Gly	Arg	Gln	Glu	Thr	Leu	Phe	Arg	185	190	195
Cys	Ile	Arg	Ser	Met	Pro	Ser	His	Pro	Asp	Arg	Ala	Tyr	Asn	Ser	200	205	210
Cys	Tyr	Ser	Ala	Gly	Val	Phe	His	Leu	His	Gln	Gly	Asp	Ile	Leu	215	220	225
Ser	Val	Ile	Ile	Pro	Arg	Ala	Arg	Ala	Lys	Leu	Asn	Leu	Ser	Pro	230	235	240
His	Gly	Thr	Phe	Leu	Gly	Phe	Val	Lys	Leu						245	250	

<210> 13

<211> 1239

<212> DNA

<213> Homo sapien

<400> 13

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ctgtggacag caccctaagc aatgtgcata cttctgtgag aacaagctca 200
ggagcccagt gaaccttcca ccagagctca ggagacagcg gagtggagaa 250
gttgaaaaca attcagacaa ctcggaaggg taccaaggat tggagcacag 300
aggctcagaa gcaagtccag ctctcccggg gctgaagctg agtgcagatc 350
agggtggcct ggtctacagc acgctggggc tctgcctgtg tgccgtcctc 400
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<210> 14  
 <211> 246  
 <212> PRT  
 <213> Homo sapien

<400> 14  
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 20 25 30  
 Lys Phe Tyr Asp His Leu Leu Arg Asp Cys Ile Ser Cys Ala Ser  
 35 40 45  
 Ile Cys Gly Gln His Pro Lys Gln Cys Ala Tyr Phe Cys Glu Asn  
 50 55 60  
 Lys Leu Arg Ser Pro Val Asn Leu Pro Pro Glu Leu Arg Arg Gln  
 65 70 75  
 Arg Ser Gly Glu Val Glu Asn Asn Ser Asp Asn Ser Gly Arg Tyr  
 80 85 90  
 Gln Gly Leu Glu His Arg Gly Ser Glu Ala Ser Pro Ala Leu Pro  
 95 100 105  
 Gly Leu Lys Leu Ser Ala Asp Gln Val Ala Leu Val Tyr Ser Thr  
 110 115 120

Leu	Gly	Leu	Cys	Leu	Cys	Ala	Val	Leu	Cys	Cys	Phe	Leu	Val	Ala	
				125					130					135	
Val	Ala	Cys	Phe	Leu	Lys	Lys	Arg	Gly	Asp	Pro	Cys	Ser	Cys	Gln	
				140					145					150	
Pro	Arg	Ser	Arg	Pro	Arg	Gln	Ser	Pro	Ala	Lys	Ser	Ser	Gln	Asp	
				155					160					165	
His	Ala	Met	Glu	Ala	Gly	Ser	Pro	Val	Ser	Thr	Ser	Pro	Glu	Pro	
				170					175					180	
Val	Glu	Thr	Cys	Ser	Phe	Cys	Phe	Pro	Glu	Cys	Arg	Ala	Pro	Thr	
				185					190					195	
Gln	Glu	Ser	Ala	Val	Thr	Pro	Gly	Thr	Pro	Asp	Pro	Thr	Cys	Ala	
				200					205					210	
Gly	Arg	Trp	Gly	Cys	His	Thr	Arg	Thr	Thr	Val	Leu	Gln	Pro	Cys	
				215					220					225	
Pro	His	Ile	Pro	Asp	Ser	Gly	Leu	Gly	Ile	Val	Cys	Val	Pro	Ala	
				230					235					240	
Gln	Glu	Gly	Gly	Pro	Gly										
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<210> 15  
 <211> 595  
 <212> DNA  
 <213> Homo sapien

<400> 15  
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 cactgcgtgg cctgcgggct cctgcgcacg ccgcggccga aaccggccgg 150  
 ggccagcagc cctgcgcccc ggacggcgct gcagccgcag gagtcggtgg 200  
 gcgcgggggc cggcgaggcg gcgctgcccc tgcccgggct gctctttggc 250  
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 tctggtgagc tggaggcggc gacagcggcg gcttcgcggc gcgtcctccg 350  
 cagaggcccc cgacggagac aaggacgccc cagagcccct ggacaaggtc 400  
 atcattctgt ctccgggaat ctctgatgcc acagctcctg cctggcctcc 450  
 tcttggggaa gaccaggaa ccaccccacc tggccacagt gtccctgtgc 500  
 cagccacaga gctgggctcc actgaactgg tgaccaccaa gacggccggc 550  
 cctgagcaac aatagcaggg agccggcagg aggtggcccc tgccc 595

<210> 16  
 <211> 184  
 <212> PRT  
 <213> Homo sapien

<400> 16

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 Pro Thr Pro Cys Val Pro Ala Glu Cys Phe Asp Leu Leu Val Arg  
 20 25 30  
 His Cys Val Ala Cys Gly Leu Leu Arg Thr Pro Arg Pro Lys Pro  
 35 40 45  
 Ala Gly Ala Ser Ser Pro Ala Pro Arg Thr Ala Leu Gln Pro Gln  
 50 55 60  
 Glu Ser Val Gly Ala Gly Ala Gly Glu Ala Ala Leu Pro Leu Pro  
 65 70 75  
 Gly Leu Leu Phe Gly Ala Pro Ala Leu Leu Gly Leu Ala Leu Val  
 80 85 90  
 Leu Ala Leu Val Leu Val Gly Leu Val Ser Trp Arg Arg Arg Gln  
 95 100 105  
 Arg Arg Leu Arg Gly Ala Ser Ser Ala Glu Ala Pro Asp Gly Asp  
 110 115 120  
 Lys Asp Ala Pro Glu Pro Leu Asp Lys Val Ile Ile Leu Ser Pro  
 125 130 135  
 Gly Ile Ser Asp Ala Thr Ala Pro Ala Trp Pro Pro Pro Gly Glu  
 140 145 150  
 Asp Pro Gly Thr Thr Pro Pro Gly His Ser Val Pro Val Pro Ala  
 155 160 165  
 Thr Glu Leu Gly Ser Thr Glu Leu Val Thr Thr Lys Thr Ala Gly  
 170 175 180  
 Pro Glu Gln Gln

<210> 17  
 <211> 265  
 <212> PRT  
 <213> Homo sapien

<400> 17  
 Met Ser Gly Leu Gly Arg Ser Arg Arg Gly Gly Arg Ser Arg Val  
 1 5 10 15  
 Asp Gln Glu Glu Arg Phe Pro Gln Gly Leu Trp Thr Gly Val Ala  
 20 25 30  
 Met Arg Ser Cys Pro Glu Glu Gln Tyr Trp Asp Pro Leu Leu Gly  
 35 40 45  
 Thr Cys Met Ser Cys Lys Thr Ile Cys Asn His Gln Ser Gln Arg  
 50 55 60  
 Thr Cys Ala Ala Phe Cys Arg Ser Leu Ser Cys Arg Lys Glu Gln  
 65 70 75  
 Gly Lys Phe Tyr Asp His Leu Leu Arg Asp Cys Ile Ser Cys Ala  
 80 85 90

Ser	Ile	Cys	Gly	Gln	His	Pro	Lys	Gln	Cys	Ala	Tyr	Phe	Cys	Glu	95	100	105
Asn	Lys	Leu	Arg	Ser	Pro	Val	Asn	Leu	Pro	Pro	Glu	Leu	Arg	Arg	110	115	120
Gln	Arg	Ser	Gly	Glu	Val	Glu	Asn	Asn	Ser	Asp	Asn	Ser	Gly	Arg	125	130	135
Tyr	Gln	Gly	Leu	Glu	His	Arg	Gly	Ser	Glu	Ala	Ser	Pro	Ala	Leu	140	145	150
Pro	Gly	Leu	Lys	Leu	Ser	Ala	Asp	Gln	Val	Ala	Leu	Val	Tyr	Ser	155	160	165
Thr	Leu	Gly	Leu	Cys	Leu	Cys	Ala	Val	Leu	Cys	Cys	Phe	Leu	Val	170	175	180
Ala	Val	Ala	Cys	Phe	Leu	Lys	Lys	Arg	Gly	Asp	Pro	Cys	Ser	Cys	185	190	195
Gln	Pro	Arg	Ser	Arg	Pro	Arg	Gln	Ser	Pro	Ala	Lys	Ser	Ser	Gln	200	205	210
Asp	His	Ala	Met	Glu	Ala	Gly	Ser	Pro	Val	Ser	Thr	Ser	Pro	Glu	215	220	225
Pro	Val	Glu	Thr	Cys	Ser	Phe	Cys	Phe	Pro	Glu	Cys	Arg	Ala	Pro	230	235	240
Thr	Gln	Glu	Ser	Ala	Val	Thr	Pro	Gly	Thr	Pro	Asp	Pro	Thr	Cys	245	250	255
Ala	Gly	Arg	Thr	Ala	Pro	Pro	Arg	Glu	Gly						260	265	